



RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

10/534966

Source:

PCR

Date Processed by STIC:

5/23/5

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA-22313-1450.
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):
U.S. Patent and Trademark Office, 220 20th Street S., Customer Window, Mail Stop Sequence, Crystal Plaza Two, Lobby, Room 1B03, Arlington, VA 22202

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER

10/534966

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>.<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>.<223> section to be missing from amino acid sequence(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>.<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>.<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO X (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS (Do not insert any subheadings under this heading!)
 (ii) SEQUENCE DESCRIPTION SEQ ID NO X (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES" response to include the skipped sequences.
- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 <210> sequence id number
 <400> sequence id number
 000
- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>.<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are Unknown, Artificial Sequence, or scientific name (Genus/species). <220>.<223> section is required when <213> response is Unknown or Artificial Sequence.
- 11 ✓ Use of <220> Sequence(s) missing the <220> "feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n/Xaa "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid.

AMC - Biotechnology Systems Branch - 09/09/2003

Best Available Copy



PCT

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/534,966

DATE: 05/23/2005

TIME: 13:51:42

Input Set : A:\8027-4-32652A.ST25.txt
 Output Set: N:\CRF4\05232005\J534966.raw

Does Not Comply
 Corrected Diskette Needed

P.1

3 <110> APPLICANT: Auer, Manfred
 4 Meisner, Nicole-Claudia
 5 Seifert, Jan-Marcus
 7 <120> TITLE OF INVENTION: Labeling methodology comprising oligopeptides
 9 <130> FILE REFERENCE: 4-32652A/NFI 8027
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/534,966
 C--> 11 <141> CURRENT FILING DATE: 2005-05-16
 11 <160> NUMBER OF SEQ ID NOS: 3
 13 <170> SOFTWARE: PatentIn version 3.1
 15 <210> SEQ ID NO: 1
 16 <211> LENGTH: 10
 17 <212> TYPE: PRT
 18 <213> ORGANISM: artificial
 20 <220> FEATURE:
 21 <223> OTHER INFORMATION: completely artificial
 23 <400> SEQUENCE: 1
 25 Cys Gly Lys Gly His His His His His His
 26 1 5 10
 29 <210> SEQ ID NO: 2
 30 <211> LENGTH: 32
 31 <212> TYPE: PRT
 32 <213> ORGANISM: Homo Sapiens
 34 <400> SEQUENCE: 2
 36 Cys Ser Asn Leu Ser Thr Cys Val Leu Gly Lys Leu Ser Gln Glu Leu
 37 1 5 10 15
 40 His Lys Leu Gln Thr Thr Pro Arg Thr Asn Thr Gly Ser Gly Thr Pro
 41 20 25 30
 44 <210> SEQ ID NO: 3
 45 <211> LENGTH: 110
 46 <212> TYPE: PRT
 47 <213> ORGANISM: Homo Sapiens
 49 <400> SEQUENCE: 3
 51 Asn Phe Pro Asn Ser Thr Asn Leu Pro Arg Asn Pro Ser Met Ala Asp
 52 1 5 10 15
 55 Tyr Glu Ala Arg Ile Phe Thr Phe Gly Thr Trp Ile Tyr Ser Val Asn
 56 20 25 30
 59 Lys Glu Gln Leu Ala Arg Ala Gly Phe Tyr Ala Leu Gly Glu Gly Asp
 60 35 40 45
 63 Lys Val Lys Cys Phe His Cys Gly Gly Gly Leu Thr Asp Trp Lys Pro
 64 50 55 60
 67 Ser Glu Asp Pro Trp Glu Gln His Ala Lys Trp Tyr Pro Gly Cys Lys
 68 65 70 75 80
 71 Tyr Leu Leu Glu Gln Lys Gly Gln Glu Tyr Ile Asn Asn Ile His Leu

INVALID RESPONSE
 MANDATORY. PLEASE
 EXPLAIN SOURCE OF
 GENETIC MATERIAL IN
 <223> SECTION. SEE
 ITEM # 11 IN ERROR
 SUMMARY SHEET.

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/534,966

DATE: 05/23/2005

TIME: 13:51:42

Input Set : A:\8027-4-32652A.ST25.txt

Output Set: N:\CRF4\05232005\J534966.raw

72				85				90				95		
75	Thr	His	Ser	Leu	Glu	Glu	Cys	Leu	Val	Arg	Thr	Thr	Glu	Lys
76				100					105					110